

Figure 1

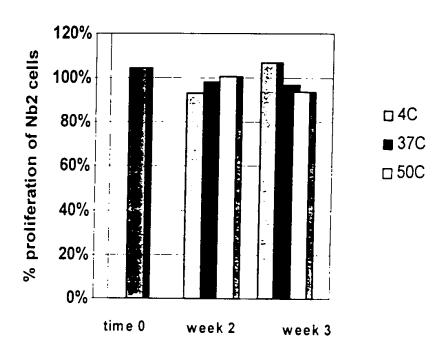


Figure 2

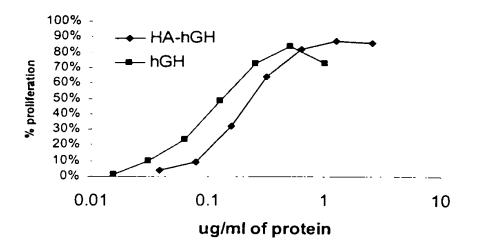


Figure 3A

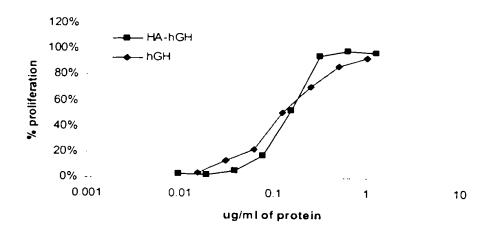


Figure 3B

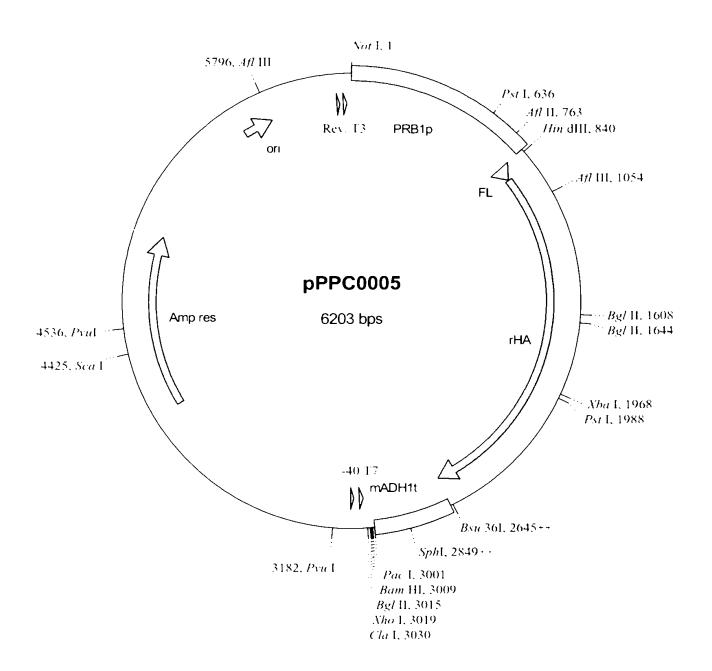


Figure 4

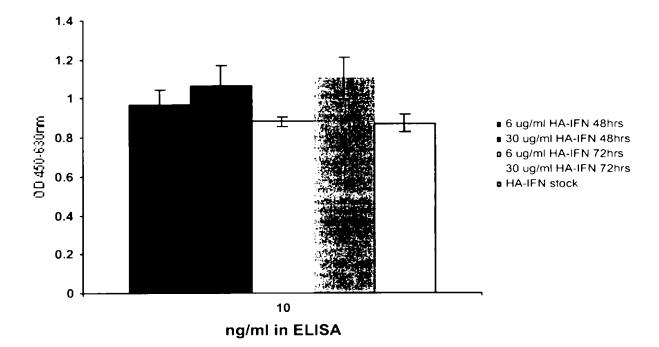
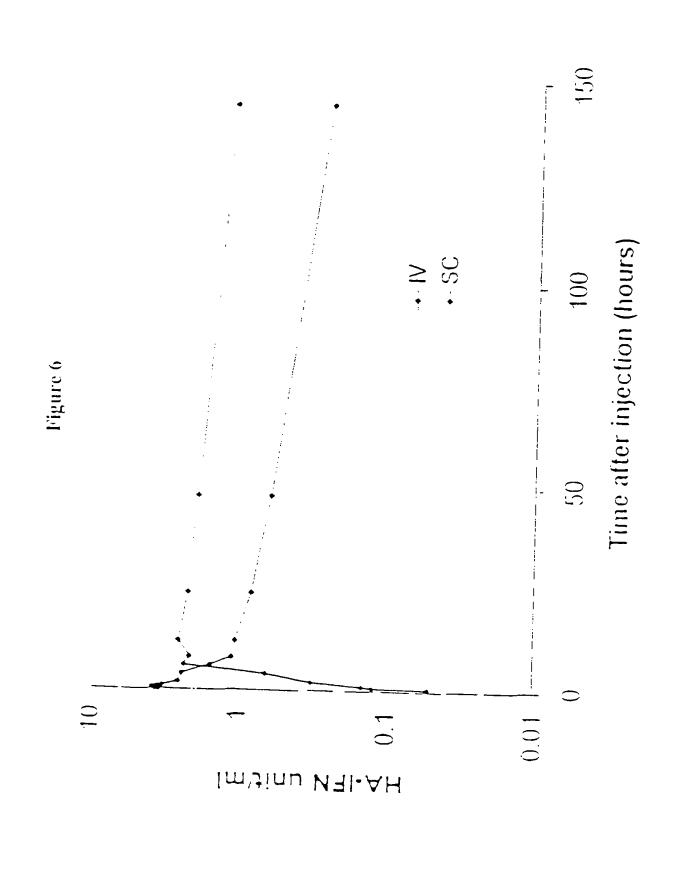
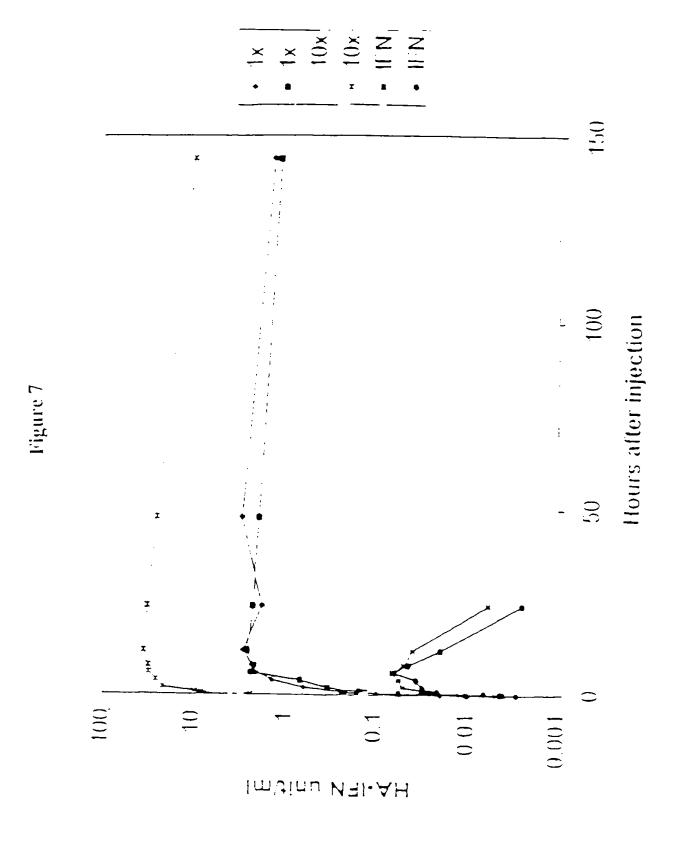


Figure 5



ΔPX (00.833,117) Got 25 Crarg A. Reservet al Arty. Docket, 6832 (10) 8300



APN 09.833444 (19.20) Crary A. Rosen et al. Alty. Docket (68.2004) 8.66

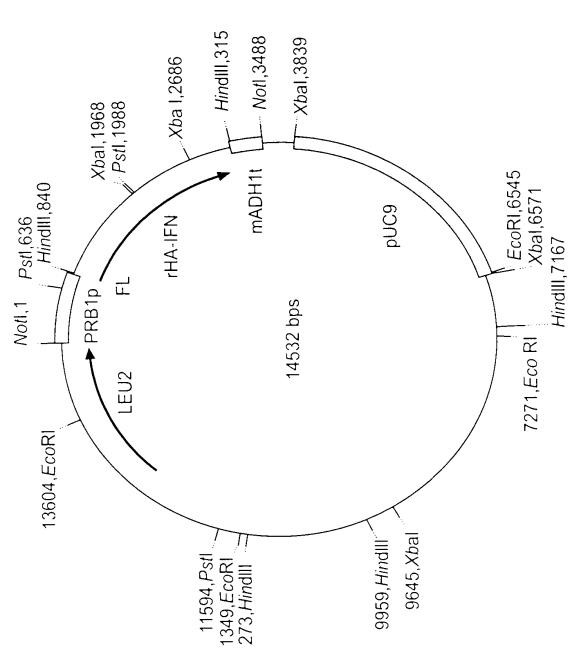


FIG. 8

## Figure 9

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		v	'I	VII	
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### Figure 10

#### a. Randomisation of Loop IV.

IV

IV

**X** represents the mutation of the natural amino acid to any other amino acid. One, more or all of the amino acids can be changed in this manner. This figure indicates all the residues have been changed.

#### b. Insertion (or replacement) of Randomised sequence into Loop IV.

(X)<sub>±</sub>

IV

The insertion can be at any point on the loop and a length where n would typically be  $\ell$ ,  $\ell$ ,  $\ell$ ,  $\ell$  or  $\ell$ .

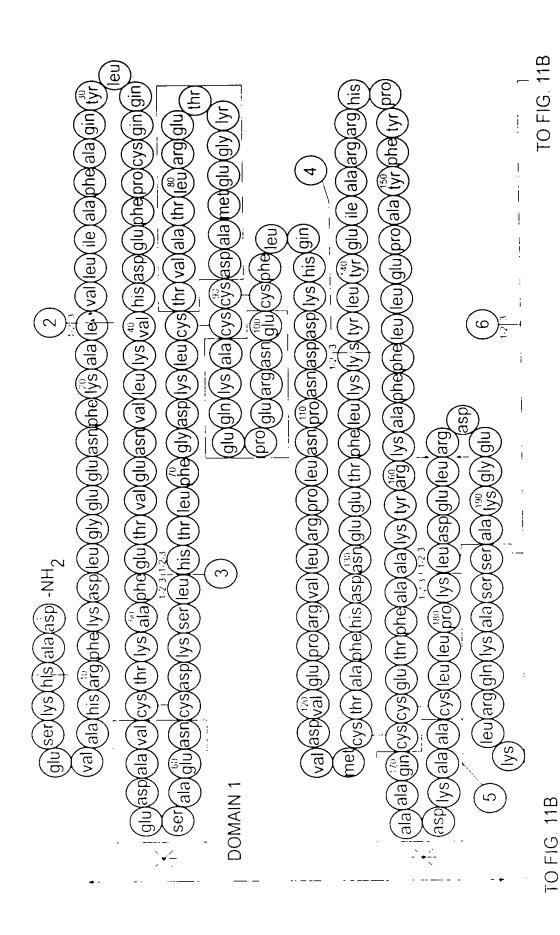


FIG. 11A

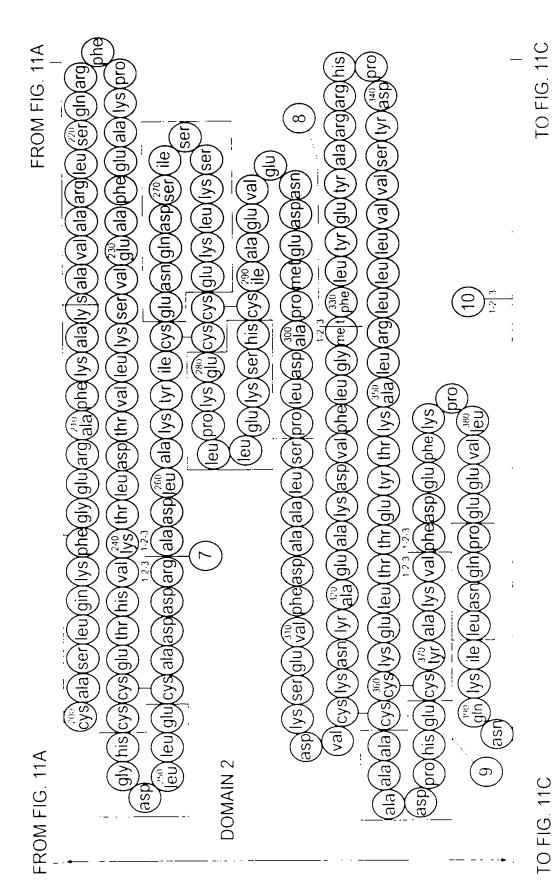
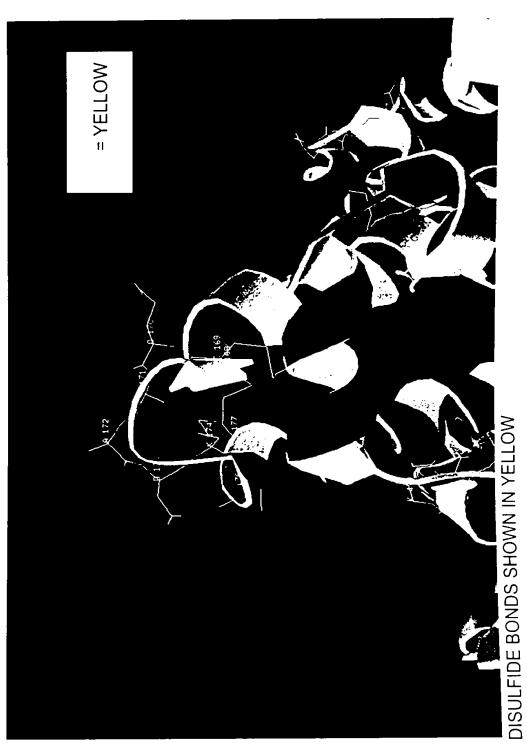


FIG. 11B

FROM FIG. 11B

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FIG. 11C



**FIG. 12:** LOOP IV GLU170-A176

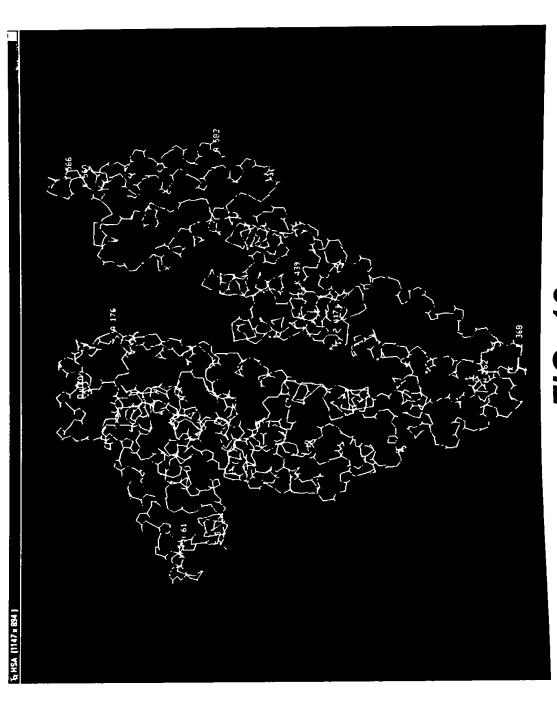
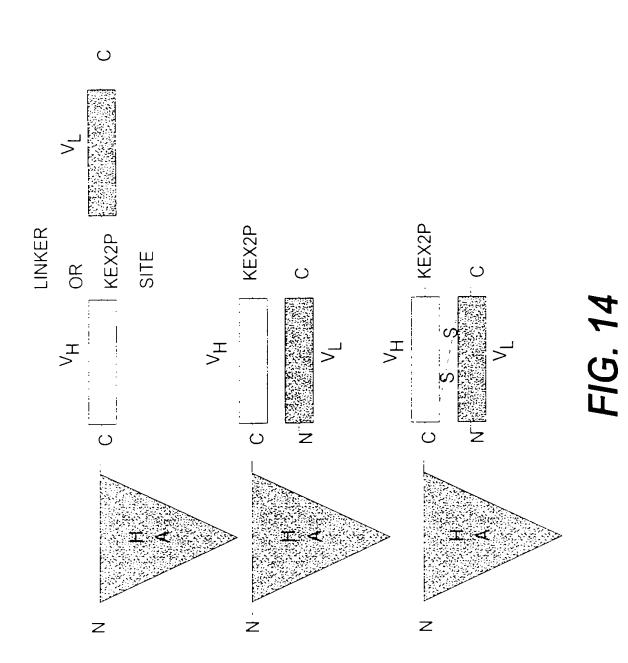


FIG. 13 TERTIARY STRUCTURE OF HA



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# Figure 15B

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Figure 15C

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Figure 15D

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